

10/586045

WO 2005/068658

PCT/KR2004/003517

JAP20 Rec'd PCT/PTO 14 JUL 2006

<110> Korea Research Institute of Bioscience and Biotechnology  
<120> Rapid screening method of translational fusion partners for  
producing recombinant proteins and translational fusion partners  
screened therefrom  
<150> KR10-2004-0003957  
<151> 2004-01-19  
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20 25 30  
Ala Asp Leu Ser Ser Ile Thr Ser Val Ser Ser Ala Ser Ala Ser Ala  
35 40 45  
Thr Ala Ser Asp Ser Leu Ser Ser Ser Asp Gly Thr Val Tyr Leu Pro  
50 55 60  
Ser Thr Thr Ile Ser Gly Asp Leu Thr Val Thr Gly Lys Val Ile Ala  
65 70 75 80  
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85 90 95  
Gly Glu Lys Tyr Val Phe Ser Ser Asp  
100 105

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aatcgttta acaaattcca agctgtgtc gcttggccc tactctctcg cggcgtctc  
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gccgtggaaag tc当地gtgccc当地 ggtaaggttg actttacttg acggtgaaaa atacgtttc  
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 20 25 30

Leu Thr Pro Thr Gly Ser Ile Ser Cys Gly Ala Ala Glu Tyr Thr Thr  
 35 40 45

Thr Phe Gly Ile Ala Val Gln Ala Ile Thr Ser Ser Lys Ala Lys Arg  
 50 55 60

Asp Val Ile Ser Gln Ile Gly Asp Gly Gln Val Gln Ala Thr Ser Ala  
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tccacacctt aa ccccaacccgg ctccatctct tttttttttt ccgaatacac taccaccc 480  
ggttttgtctg ttcaagctat taccttttca aaagctaaaga gagacgttat ctctcaaatt 540  
ggtgacggtc aagtccaaagc cactttgtct gctactgttc aagccaccggaa tagtcaagcc 600  
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Lys 11e  
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20 25 30  
Ala Asp Leu Ser Ser Ile Thr Ser Val Ser Ser Ala Ser Ala Ser Ala  
35 40 45  
Thr Ala Ser Asp Ser Leu Ser Ser Ser Asp Gly Thr Val Tyr Leu Pro  
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Ser Thr Thr Ile Ser Gly Asp  
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acttccgtct cgtcagctag tgcaagtgcc accgcctccg actcacattc ttccagtgac 180  
ggtaaccgttt atttgcacatc cacaacaattt agcgggtgatc tcacagttac tggtaaagta 240  
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<212> DNA  
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<223> JH97(Sfi-HSA-forward primer)

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<212> DNA  
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29

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<212> DNA  
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<223> JH99(Sfi-INV-forward primer)

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37

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37

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26

<210> 18  
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22

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22

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<210> 24  
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<212> DNA  
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47

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25

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Leu Thr Pro Thr Gly Ser Ile Ser Cys Gly Ala Ala Glu Tyr Thr Thr  
 35 40 45

Thr Phe Gly Ile Ala Val Gln Ala Ile Thr Ser Ser Lys Ala Lys Arg  
 50 55 60

Asp Val Ile Ser Gln Ile Gly Asp Gly Gln Val Gln Ala Thr Ser Ala  
 65 70 75 80

Ala Thr Ala Gln Ala Thr Asp Ser Gln Ala Gln Ala Thr Thr Thr Ala  
 85 90 95

Thr Pro Thr Ser Ser Glu Lys Ile Ser Ser Ser Ala Ser Lys Thr Ser  
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tgtggtgctg ccgaatacac taccaccttt ggtattgctg ttcaagctat taccctttca 180  
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Leu Thr Pro Thr Gly Ser Ile Ser Cys Gly Ala Ala Glu Tyr Thr Thr  
 35 40 45

Thr Phe Gly Ile Ala Val Gln Ala Ile Thr Ser Ser Lys Ala Lys Arg  
 50 55 60

Asp Val Ile Ser Gln Ile Gly Asp Gly Gln Val Gln Ala Thr Ser Ala  
 65 70 75 80

Ala Thr Ala Gln Ala Thr Asp Ser Gln Ala Gln Ala Thr Thr Thr Ala

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Thr Pro Thr Ser Ser Glu Lys Ile Ser Ser Ser Ala Ser Lys Thr Ser  
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Thr Asn Ala Thr Ser Ser Ser Cys Ala Thr Pro Ser Leu Lys Asp Ser  
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tgtggtgctg	ccgaatacac	taccacctt	ggtattgttg	ttcaaggtat	tacccatcttca	180
aaagctaaga	gagacgttat	ctctcaaatt	ggtgacggtc	aagtccaagc	cacttctgt	240
gtctactgctc	aaggccaccga	tagtcaagcc	caagctacta	ctaccgtac	cccaaccaggc	300
tccgaaaaga	tctcttcctc	tgcatactaaa	acatctacta	atgccacatc	atcttcttgt	360
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